FIG. 15

Influence of Y1-scFv on platelets agglutination in washed platelets

% Agglutination

TM1-scFv 25 ug/ml · Y1-scFv 25ug/ml · Y1-scFv 12.5 ug/ml · Y1-scFv 4 ug/ml · Y17-scFv 25ug/ml · Y16-scFv 38 ug/ml

concentration

REPLACEMENT SHEET

FIG. 25

α-CD162 and Y1:
Comparison between cells
from AML patient and normal
blood
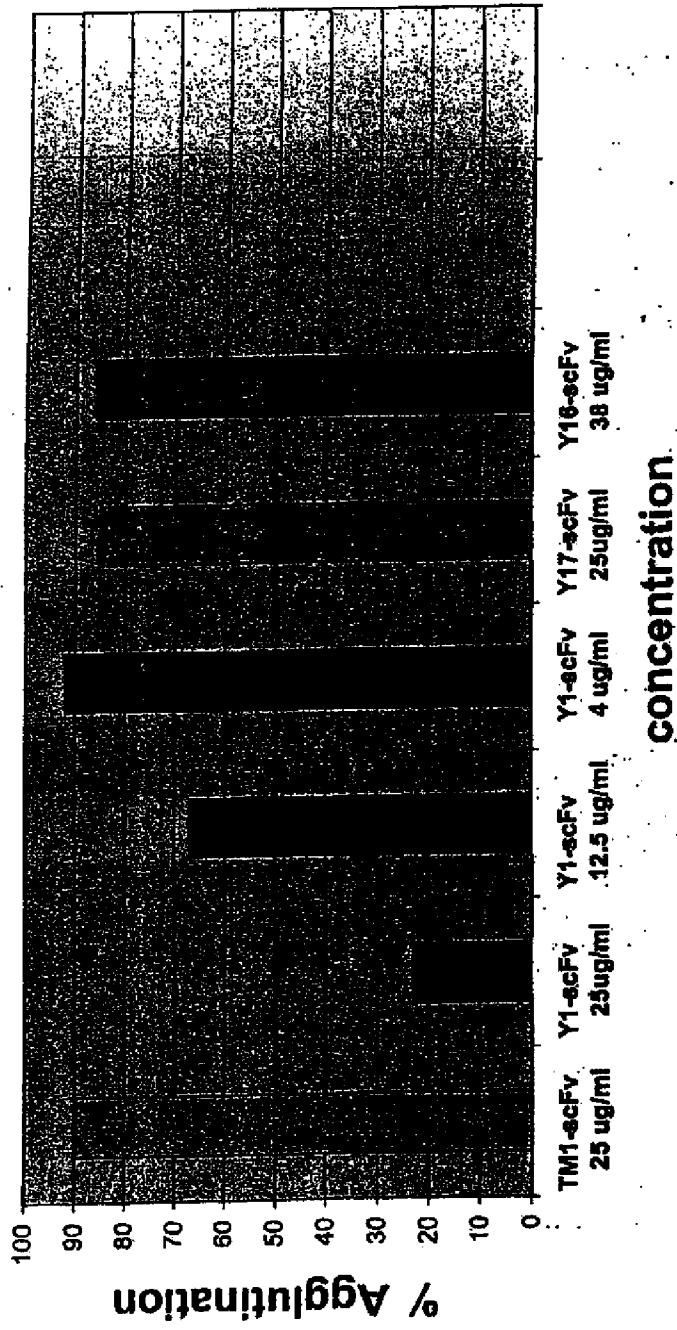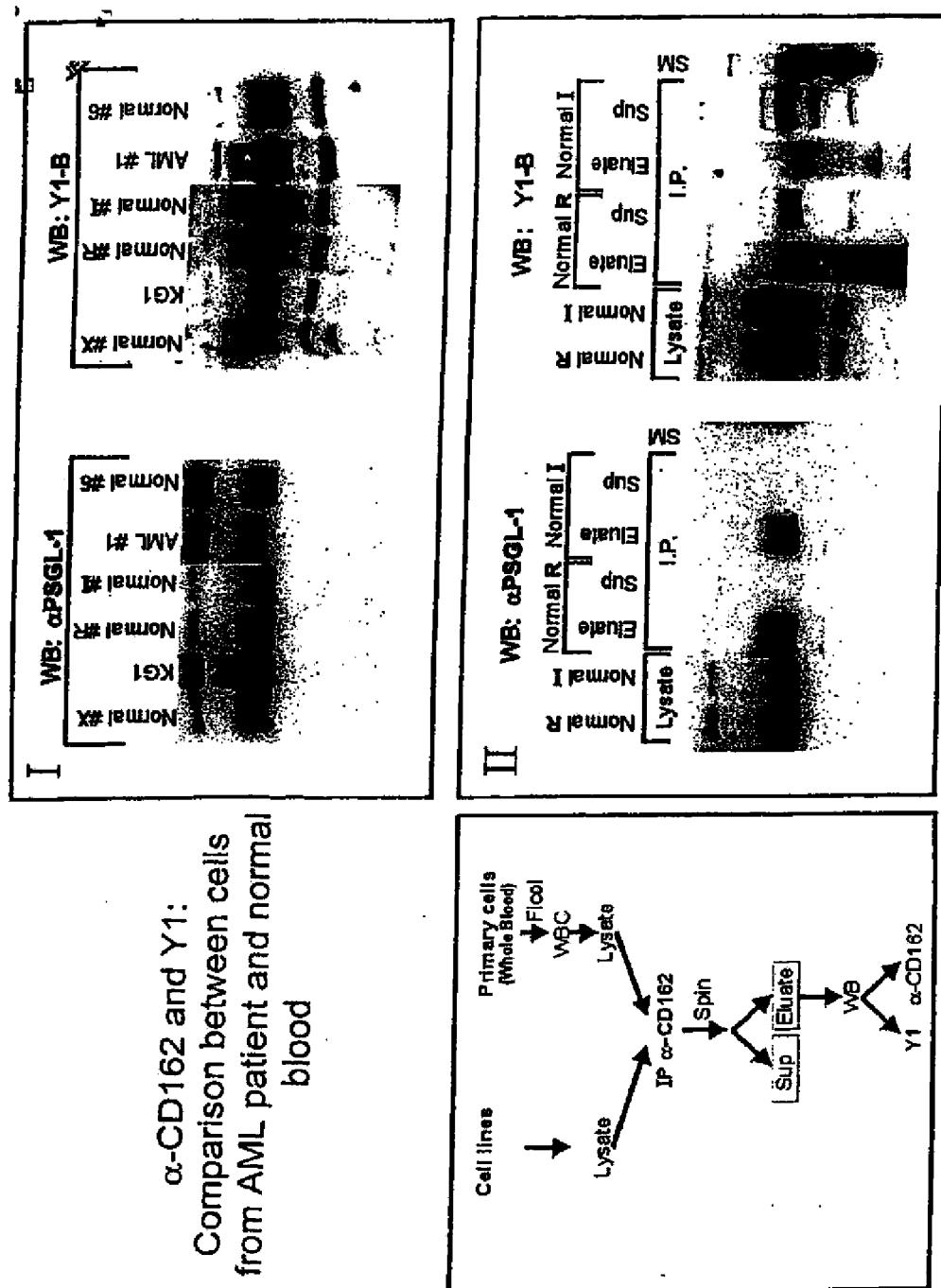
REPLACEMENT SHEET

FIG. 50

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        10          20          30          40          50          60
ActTaTTActc gCGGCCCAGC CgGCCxxxxxC CgAGGTGCAG CTGGTGGAGT CTGGGGGAGG
 L  L  L  x  A  Q  P  A  x  x  x    E  V  Q  L  V  E  S  G  G

        70          80          90          100         110         120
CTTGGTACAG CCTGGGGGGT CCCTGAGACT CTCCTGTGCA GCCTCTGGAT TCACCTTTAG
 L  V  Q  P  G  G  S  L  R  L  S  C  A  A  S  G  F  T  F  S

        130         140         150         160         170         180
CAGCTATGCC ATGAGCTGGG TCCGCCAGGC TCCAGGGAAG GGGCTGGAGT GGGTCTCAGC
 S  Y  A  M  S  W  V  R  Q  A  P  G  K  G  L  E  W  V  S  A

        190         200         210         220         230         240
TATTAGTGGT AGTGGTGGTA GCACATACTA CGCAGACTCC GTGAAGGGCC GGTTCACCAT
 I  S  G  S  G  G  S  T  Y  Y  A  D  S  V  K  G  R  F  T  I

        250         260         270         280         290         300
CTCCAGAGAC AATTCCAAGA ACACGCTGTA TCTGCAAATG AACAGCCTGA GAGCCGAGGA
 S  R  D  N  S  K  N  T  L  Y  L  Q  M  N  S  L  R  A  E  D

        310         320         330          340        350         360
CACGGCCGTG TATTACTGTG CAAGAxxxx-GCAGxxxxxxxxxxxGxxxT GGGGCCAAGG
 T  A  V  Y  Y  C  A  R  L  G  Q  S  I  K  R  S  W  G  Q  G

        370         380         390         400         410         420
TACCCTGGTC ACCGTGTCGA GAGGTGGAGG CGGTTCAGGC GGAGGTGGCT CTGGCGGTGG
 T  L  V  T  V  S  R  G  G  G  S  G  G  G  S  G  G  G

        430         440         450         460         470         480
CGGATGGTCT GAGCTGACTC AGGACCCTGC TGTGTCTGTG GCCTTGGGAC AGACAGTCAG
 G  S  S  E  L  T  Q  D  P  A  V  S  V  A  L  G  Q  T  V  R

        490         500         510         520         530         540
GATCACATGC CAAGGAGACA GCCTCAGAAG CTATTATGCA AGCTGGTACC AGCAGAAGCC
 I  T  C  Q  G  D  S  L  R  S  Y  Y  A  S  W  Y  Q  Q  K  P

        550         560         570         580         590         600
AGGACAGGCC CCTGTACTTG TCATCTATGG TAAAAACAAC CGGCCCTCAG GGATCCCAGA
 G  Q  A  P  V  L  V  I  Y  G  K  N  N  R  P  S  G  I  P  D

        610         620         630         640         650         660
CCGATTCTCT GGCTCCAGCT CAGGAAACAC AGCTTCCTTG ACCATCACTG GGGCTCAGGC
 R  F  S  G  S  S  S  G  N  T  A  S  L  T  I  T  G  A  Q  A

        670         680         690         700         710         720
GGAAGATGAG GCTGACTATT ACTGTAACTC CCGGGACAGC AGTGGTAACC ATGTGGTATT
 E  D  E  A  D  Y  Y  C  N  S  R  D  S  S  G  N  H  V  V  F

        730         740         750         760         770         780
CGGCGGAGGG ACCAAGCTGA CCGTCCTAGG TGCGGCCGCA GAACAAAAAC TCATCTCAGA
 G  G  G  T  K  L  T  V  L  G  A  A  A  E  Q  K  L  I  S  E

        790         800         810         820         830         840
AGAGGAtCTG AAtGGGGCCG CAxxxxACTG TtGAATTTTT TAAGTtAACC T
 E  D  L  N  G  A  A  *  N  C  *  I  F  *  V  N
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